Document made available under the Patent Cooperation Treaty (PCT)

International application number: PCT/EP05/000562

International filing date: 18 January 2005 (18.01.2005)

Document type:

Certified copy of priority document

Document details:

Country/Office: EP

04100206.4

Number:

Filing date: 22 January 2004 (22.01.2004)

Date of receipt at the International Bureau: 24 February 2005 (24.02.2005)

Remark:

Priority document submitted or transmitted to the International Bureau in

compliance with Rule 17.1(a) or (b)



World Intellectual Property Organization (WIPO) - Geneva, Switzerland Organisation Mondiale de la Propriété Intellectuelle (OMPI) - Genève, Suisse



Europäisches Patentamt

European **Patent Office** Office européen des brevets

10.01.05

Bescheinigung

Certificate

Attestation

Die angehefteten Unterlagen stimmen mit der ursprünglich eingereichten Fassung der auf dem nächsten Blatt bezeichneten europäischen Patentanmeldung überein.

The attached documents are exact copies of the European patent application conformes à la version described on the following initialement déposée de page, as originally filed.

Les documents fixés à cette attestation sont initialement déposée de la demande de brevet européen spécifiée à la page suivante.

Patentanmeldung Nr.

Patent application No. Demande de brevet n°

04100206.4

Der Präsident des Europäischen Patentamts; Im Auftrag

For the President of the European Patent Office

Le Président de l'Office européen des brevets

R C van Dijk



Europäisches Patentamt

European Patent Office Office européen des brevets

PCT/EP200 5 / 0 0 0 5 6 2

10.01.05

Anmeldung Nr:

Application no.: 04

04100206.4

Demande no:

Anmeldetag:

Date of filing: 22.01.04

Date de dépôt:

Anmelder/Applicant(s)/Demandeur(s):

Akzo Nobel N.V. Velperweg 76 6824 BM Arnhem PAYS-BAS

Bezeichnung der Erfindung/Title of the invention/Titre de l'invention: (Falls die Bezeichnung der Erfindung nicht angegeben ist, siehe Beschreibung. If no title is shown please refer to the description. Si aucun titre n'est indiqué se referer à la description.)

LAWSONIA INTRACELLULARIS 74 kD SUBUNIT VACCINE

In Anspruch genommene Prioriät(en) / Priority(ies) claimed /Priorité(s) revendiquée(s)
Staat/Tag/Aktenzeichen/State/Date/File no./Pays/Date/Numéro de dépôt:

Internationale Patentklassifikation/International Patent Classification/Classification internationale des brevets:

C07K14/00

Am Anmeldetag benannte Vertragstaaten/Contracting states designated at date of filing/Etats contractants désignées lors du dépôt:

AT BE BG CH CY CZ DE DK EE ES FI FR GB GR HU IE IT LU MC NL PT RO SE SI SK TR LI

Lawsonia intracellularis 74 kD subunit vaccine.

The present invention relates i.a. to nucleic acids encoding novel Lawsonia intracellularis proteins, to DNA fragments, recombinant DNA molecules and live recombinant carriers comprising these sequences, to host cells comprising such nucleic acids, DNA fragments, recombinant DNA molecules and live recombinant carriers, to proteins encoded by these nucleotide sequences and to their use for the manufacturing of vaccines, to vaccines for combating Lawsonia intracellularis infections and methods for the preparation thereof and to diagnostic tests for the detection of Lawsonia intracellularis antigens and for the detection of antibodies against Lawsonia intracellularis.

Porcine proliferative enteropathy (PPE or PE) has become an important disease of the modern pig industry world-wide. The disease affects 15% to 50% of the growing herds and up to 30% of the individual animals in established problem herds. Today annual economical losses have been estimated US\$ 5-10 in extra feed and facility time costs per affected pig. PPE is a group of chronic and acute conditions of widely differing clinical signs (death, pale and anaemic animals, watery, dark or bright red diarrhoea, depression, reduced appetite and reluctance to move, retarded growth and increased FCR). However there are two consistent features. The first, a pathological change only visible at necropsy, is a thickening of the small intestine and colon mucosa. The second is the occurrence of intracytoplasmatic small-curved bacteria in the enterocytes of the affected intestine. These bacteria have now been established as the etiological agent of PPE and have been name Lawsonia intracellularis.

25

30

5

10

15

20

Over the years Lawsonia intracellularis has been found to affect a large group of animals including monkeys, rabbits, ferrets, hamsters, fox, horses, and other animals as diverse as ostrich and emoe. Lawsonia intracellularis is a gram-negative, flagellated bacterium that multiplies in eukaryotic enterocytes only and no cell-free culture has been described. In order to persist and multiply in the cell Lawsonia intracellularis must penetrate dividing crypt cells. The bacterium associates with the cell membrane and quickly enters the enterocyte via an entry vacuole. This then rapidly breaks down (within 3 hours) and the bacteria flourish and multiply freely in the cytoplasm. The mechanisms by which the bacteria cause infected cells to fail to

mature, continue to undergo mitosis and form hypoplastic crypt cells is not yet understood.

The current understanding of Lawsonia intracellularis infection, treatment and control of the disease has been hampered by the fact that Lawsonia intracellularis can not be cultivated in cell-free media. Although there are reports of successful co-culturing Lawsonia intracellularis in rat enterocytes this has not lead to the development of inactivated vaccines for combating Lawsonia intracellularis, although there clearly is a need for such vaccines.

10

25

It is an objective of the present invention to provide a vaccine for combating Lawsonia intracellularis infection.

It was surprisingly found now, that Lawsonia intracellularis produces a novel protein that is capable of inducing protective immunity against Lawsonia intracellularis.

The novel protein will be referred to as the 74 kD protein.

The amino acid sequence of the novel protein is presented in sequence identifier SEQ ID NO: 2. The gene encoding this protein has been sequenced and its nucleic acid sequence is shown in sequence identifier SEQ ID NO: 1. The gene will also be referred to in the Examples as "gene 5293".

It is well-known in the art, that many different nucleic acid sequences can encode one and the same protein. This phenomenon is commonly known as wobble in the second and especially the third base of each triplet encoding an amino acid. This phenomenon can result in a heterology of about 30% for two nucleic acid sequences still encoding the same protein. Therefore, two nucleic acid sequences having a sequence homology of about 70 % can still encode one and the same protein.

Thus, one embodiment relates to nucleic acids encoding a Lawsonia intracellularis protein and to parts of that nucleic acid that encode an immunogenic fragment of that protein, wherein those nucleic acids or parts thereof have a level of homology with the nucleic acid of which the sequence is given in SEQ ID NO: 1 of at least 90 %.

Preferably, the nucleic acid encoding this Lawsonia intracellularis protein or the part of said nucleic acid has at least 92 %, preferably 94 %, more preferably 95 % and even more preferably 96% homology with the nucleic acid having the sequence given in SEQ ID NO: 1. Even more preferred is a homology level of 98 % or even 100 %.

5

The level of nucleotide homology can be determined with the computer program "BLAST 2 SEQUENCES" by selecting sub-program: "BLASTN" that can be found at www.ncbi.nlm.nih.gov/blast/bl2seq/bl2.html.

A reference for this program is Tatiana A. Tatusova, Thomas L. Madden FEMS

Microbiol. Letters 174: 247-250 (1999). Parameters used are the default parameters:

Reward for a match: +1. Penalty for a mismatch: -2. Open gap: 5. Extension gap: 2.

Gap x_dropoff: 50.

Another approach for deciding if a certain nucleic acid is or is not a nucleic acid according to the invention relates to the question if that certain nucleic acid does hybridise under stringent conditions to nucleic acids having the nucleotide sequence as depicted in SEQ ID NO: 1.

If a nucleic acid hybridises under stringent conditions to the nucleotide sequence as depicted in SEQ ID NO: 1, it is considered to be a nucleic acid according to the invention.

The definition of stringent conditions follows from the formula of Meinkoth and Wahl (1984. Hybridization of nucleic acids immobilized on solid supports. Anal. Biochem. 138: 267-284.)

 $Tm = [81.5^{\circ}C + 16.6(log M) + 0.41(\%GC) - 0.61(\%formamide) - 500/L] - 1^{\circ}C/1\%mismatch$

In this formula, M is molarity of monovalent cations; %GC is the percentage of guanosine and cytosine nucleotides in the DNA; L is the length of the hybrid in base pairs.

Stringent conditions are those conditions under which nucleic acids or fragments thereof still hybridise, if they have a mismatch of 10% at the most, to the nucleic acid having the sequence depicted in SEQ ID NO: 1.

Since the present invention discloses nucleic acids encoding novel Lawsonia 5 intracellularis proteins, it is now for the first time possible to obtain these proteins in sufficient quantities. This can e.g. be done by using expression systems to express the genes encoding the proteins.

Therefore, in a more preferred embodiment, the invention relates to DNA fragments comprising a nucleic acid according to the invention. Such DNA fragments can e.g. 10 be plasmids, into which a nucleic acid according to the invention is cloned. Such DNA fragments are e.g. useful for enhancing the amount of DNA for use as a primer, as described below.

An essential requirement for the expression of the nucleic acid is an adequate 15 promoter functionally linked to the nucleic acid, so that the nucleic acid is under the control of the promoter. It is obvious to those skilled in the art that the choice of a promoter extends to any eukaryotic, prokaryotic or viral promoter capable of directing gene transcription in cells used as host cells for protein expression.

Therefore, an even more preferred form of this embodiment relates to a recombinant 20 DNA molecule comprising a DNA fragment or a nucleic acid according to the invention that is placed under the control of a functionally linked promoter. This can be acomplished by means of e.g. standard molecular biology techniques. (Sambrook, J. and Russell, D.W., Molecular cloning: a laboratory manual, 2001. ISBN 0-87969-

577-3). 25

30

Functionally linked promoters are promoters that are capable of controlling the transcription of the nucleic acids to which they are linked.

Such a promoter can be a Lawsonia promoter e.g. the promoter involved in in vivo expression of the gene encoding the 74 kD protein, provided that that promoter is functional in the cell used for expression. It can also be a heterologous promoter. When the host cells are bacteria, useful expression control sequences which may be used include the Trp promoter and operator (Goeddel, et al., Nucl. Acids Res., 8, 4057, 1980); the lac promoter and operator (Chang, et al., Nature, 275, 615, 1978); the outer membrane protein promoter (Nakamura, K. and Inouge, M., EMBO J., 1, 771775, 1982); the bacteriophage lambda promoters and operators (Remaut, E. et al., Nucl. Acids Res., 11, 4677-4688, 1983); the α-amylase (B. subtilis) promoter and operator, termination sequences and other expression enhancement and control sequences compatible with the selected host cell.

When the host cell is yeast, useful expression control sequences include, e.g., α-mating factor. For insect cells the polyhedrin or p10 promoters of baculoviruses can be used (Smith, G.E. et al., Mol. Cell. Biol. 3, 2156-65, 1983). When the host cell is of mammalian origin illustrative useful expression control sequences include the SV-40 promoter (Berman, P.W. et al., Science, 222, 524-527, 1983) or the
metallothionein promoter (Brinster, R.L., Nature, 296, 39-42, 1982) or a heat shock promoter (Voellmy et al., Proc. Natl. Acad. Sci. USA, 82, 4949-53, 1985).

Bacterial, yeast, fungal, insect and mammalian cell expression systems are very frequently used systems. Such systems are well-known in the art and generally available, e.g. commercially through Invitrogen (www.invitrogen.com), Novagen (www.merckbiosciences.de) or Clontech Laboratories, Inc. 4030 Fabian Way, Palo Alto, California 94303-4607, USA. Next to these expression systems, parasite-based expression systems are very attractive expression systems. Such systems are e.g. described in the French Patent Application with Publication number 2 714 074, and in US NTIS Publication No US 08/043109 (Hoffman, S. and Rogers, W.: Public. Date 1 December 1993).

15

20

A still even more preferred form of this embodiment of the invention relates to Live Recombinant Carriers (LRCs) comprising a nucleic acid encoding the 74 kD protein or an immunogenic fragment thereof according to the invention, a DNA fragment according to the invention or a recombinant DNA molecule according to the invention. Such carriers are e.g. bacteria and viruses. These LRCs are microorganisms or viruses in which additional genetic information, in this case a nucleic acid encoding the 74 kD protein or an immunogenic fragment thereof according to the invention has been cloned. Animals infected with such LRCs will produce an immunogenic response not only against the immunogens of the carrier, but also against the immunogenic parts of the protein(s) for which the genetic code is additionally cloned into the LRC, e.g. the 74 kD protein.

As an example of bacterial LRCs, attenuated Salmonella strains known in the art can attractively be used.

Live recombinant carrier parasites have i.a. been described by Vermeulen, A. N. (Int. Journ. Parasitol. 28: 1121-1130 (1998))

Also, LRC viruses may be used as a way of transporting the nucleic acid into a target cell. Live recombinant carrier viruses are also called vector viruses. Viruses often used as vectors are Vaccinia viruses (Panicali et al; Proc. Natl. Acad. Sci. USA, 79: 4927 (1982), Herpesviruses (E.P.A. 0473210A2), and Retroviruses (Valerio, D. et al; in Baum, S.J., Dicke, K.A., Lotzova, E. and Pluznik, D.H. (Eds.), Experimental Haematology today - 1988. Springer Verlag, New York: pp. 92-99 (1989)).

The technique of *in vivo* homologous recombination, well-known in the art, can be used to introduce a recombinant nucleic acid into the genome of a bacterium, parasite or virus of choice, capable of inducing expression of the inserted nucleic acid according to the invention in the host animal.

15

Finally another form of this embodiment of the invention relates to a host cell comprising a nucleic acid encoding a protein according to the invention, a DNA fragment comprising such a nucleic acid or a recombinant DNA molecule comprising such a nucleic acid under the control of a functionally linked promoter. This form also 20 relates to a host cell containing a live recombinant carrier containing a nucleic acid molecule encoding a 74 kD protein or a fragment thereof according to the invention. A host cell may be a cell of bacterial origin, e.g. Escherichia coli, Bacillus subtilis and Lactobacillus species, in combination with bacteria-based plasmids as pBR322, or bacterial expression vectors as pGEX, or with bacteriophages. The host cell may also 25 be of eukaryotic origin, e.g. yeast-cells in combination with yeast-specific vector molecules, or higher eukaryotic cells like insect cells (Luckow et al; Bio-technology 6: 47-55 (1988)) in combination with vectors or recombinant baculoviruses, plant cells in combination with e.g. Ti-plasmid based vectors or plant viral vectors (Barton, K.A. et al; Cell 32: 1033 (1983), mammalian cells like Hela cells, Chinese Hamster 30 Ovary cells (CHO) or Crandell Feline Kidney-cells, also with appropriate vectors or recombinant viruses.

Another embodiment of the invention relates to the novel proteins and to immunogenic fragments thereof according to the invention.

The concept of immunogenic fragments will be defined below.

5

One form of this embodiment relates i.a. to Lawsonia intracellularis proteins that have an amino acid sequence that is at least 90 % homologous to the amino acid sequence as depicted in SEQ ID NO: 2 and to immunogenic fragments of said protein.

In a preferred form, the embodiment relates to such Lawsonia intracellularis proteins that have a sequence homology of at least 92 %, preferably 94 %, more preferably 96 % homology to the amino acid sequence as depicted in SEQ ID NO: 2 and to immunogenic fragments of such proteins.

Even more preferred is a homology level of 98 % or even 100 %.

15

25

30

The level of protein homology can be determined with the computer program "BLAST 2 SEQUENCES" by selecting sub-program: "BLASTP", that can be found at www.ncbi.nlm.nih.gov/blast/bl2seq/bl2.html.

A reference for this program is Tatiana A. Tatusova, Thomas L. Madden FEMS

Microbiol. Letters 174: 247-250 (1999). Matrix used: "blosum62". Parameters used are the default parameters:

Open gap: 11. Extension gap: 1. Gap x_dropoff: 50.

It will be understood that, for the particular proteins embraced herein, natural variations can exist between individual Lawsonia intracellularis strains. These variations may be demonstrated by (an) amino acid difference(s) in the overall sequence or by deletions, substitutions, insertions, inversions or additions of (an) amino acid(s) in said sequence. Amino acid substitutions which do not essentially alter biological and immunological activities, have been described, e.g. by Neurath et al in "The Proteins" Academic Press New York (1979). Amino acid replacements between related amino acids or replacements which have occurred frequently in evolution are, inter alia, Ser/Ala, Ser/Gly, Asp/Gly, Asp/Asn, Ile/Val (see Dayhof, M.D., Atlas of protein sequence and structure, Nat. Biomed. Res. Found., Washington D.C., 1978, vol. 5, suppl. 3). Other amino acid substitutions include Asp/Glu, Thr/Ser,

Ala/Gly, Ala/Thr, Ser/Asn, Ala/Val, Thr/Phe, Ala/Pro, Lys/Arg, Leu/Ile, Leu/Val and Ala/Glu. Based on this information, Lipman and Pearson developed a method for rapid and sensitive protein comparison (Science,227, 1435-1441, 1985) and determining the functional similarity between homologous proteins. Such amino acid substitutions of the exemplary embodiments of this invention, as well as variations having deletions and/or insertions are within the scope of the invention as long as the resulting proteins retain their immune reactivity.

This explains why Lawsonia intracellularis proteins according to the invention, when isolated from different field isolates, may have homology levels of about 90 %, while still representing the same protein with the same immunological characteristics.

Those variations in the amino acid sequence of a certain protein according to the invention that still provide a protein capable of inducing an immune response against infection with Lawsonia intracellularis or at least against the clinical manifestations of the infection are considered as "not essentially influencing the immunogenicity".

15

20

25

30

5

10

When a protein is used for e.g. vaccination purposes or for raising antibodies, it is however not necessary to use the whole protein. It is also possible to use a fragment of that protein that is capable, as such or coupled to a carrier such as e.g. KLH, of inducing an immune response against that protein, a so-called immunogenic fragment. An "immunogenic fragment" is understood to be a fragment of the full-length protein that still has retained its capability to induce an immune response in the host, i.e. comprises a B- or T-cell epitope. At this moment, a variety of techniques is available to easily identify DNA fragments encoding antigenic fragments (determinants). The method described by Geysen et al (Patent Application WO 84/03564, Patent Application WO 86/06487, US Patent NR. 4,833,092, Proc. Natl Acad. Sci. 81: 3998-4002 (1984), J. Imm. Meth. 102, 259-274 (1987), the so-called PEPSCAN method is an easy to perform, quick and well-established method for the detection of epitopes; the immunologically important regions of the protein. The method is used world-wide and as such well-known to man skilled in the art. This (empirical) method is especially suitable for the detection of B-cell epitopes. Also, given the sequence of the gene encoding any protein, computer algorithms are able to designate specific protein fragments as the immunologically important epitopes on the basis of their sequential and/or structural agreement with epitopes that are now known. The determination of these regions is based on a combination of the hydrophilicity criteria according to

Hopp and Woods (Proc. Natl. Acad. Sci. 78: 38248-3828 (1981)), and the secondary structure aspects according to Chou and Fasman (Advances in Enzymology 47: 45-148 (1987) and US Patent 4,554,101). T-cell epitopes can likewise be predicted from the sequence by computer with the aid of Berzofsky's amphiphilicity criterion (Science 235, 1059-1062 (1987) and US Patent application NTIS US 07/005,885). A condensed overview is found in: Shan Lu on common principles: Tibtech 9: 238-242 (1991), Good et al on Malaria epitopes; Science 235: 1059-1062 (1987), Lu for a review; Vaccine 10: 3-7 (1992), Berzowsky for HIV-epitopes; The FASEB Journal 5:2412-2418 (1991).

10

5

Therefore, one form of still another embodiment of the invention relates to vaccines capable of protecting pigs against *Lawsonia intracellularis* infection, that comprise a protein or an immunogenic fragment thereof, according to the invention as described above together with a pharmaceutically acceptable carrier.

15

25

30

Still another embodiment of the present invention relates to the proteins according to the invention for use in a vaccine.

Still another embodiment relates to the use of a protein according to the invention for the manufacturing of a vaccine for combating Lawsonia intracellularis infections.

One way of making a vaccine according to the invention is by biochemical purification of the proteins or immunogenic fragments thereof according to the invention from bacteria obtained through mucosal scrapings taken from the infected intestine wall. This is however a very time-consuming way of making the vaccine.

It is therefore much more convenient to use the expression products of the genes encoding the proteins or immunogenic fragments thereof according to the invention in vaccines. The nucleic acid of the gene encoding the 74 kD protein is provided by the present invention.

Such vaccines based upon the expression products of these genes can easily be made by admixing a protein according to the invention or an immunogenic fragment thereof according to the invention with a pharmaceutically acceptable carrier as described below.

Alternatively, a vaccine according to the invention can comprise live recombinant carriers as described above, capable of expressing the proteins according to the invention or immunogenic fragments thereof according to the invention. Such vaccines, e.g. based upon a Salmonella carrier or a viral carrier infecting the enteric epithelium, or e.g. the respiratory epithelium have the advantage over subunit vaccines that they better mimic the natural way of infection of Lawsonia intracellularis. Moreover, their self-propagation is an advantage since only low amounts of the recombinant carrier are necessary for immunisation.

5

10

30

Vaccines described above all contribute to active vaccination, i.e. the host's immune system is triggered by a protein according to the invention or an immunogenic fragment thereof, to make antibodies against these proteins. Alternatively, such antibodies can be raised in e.g. rabbits or can be obtained from antibody-producing cell lines as described below. Such antibodies can then be administered to the host animal. This method of vaccination, passive vaccination, is the vaccination of choice when an animal is already infected, and there is no time to allow the natural immune response to be triggered. It is also the preferred method for 20 vaccinating immune-compromised animals. Administered antibodies against Lawsonia intracellularis can in these cases bind directly to the bacteria. This has the advantage that it immediately decreases or stops Lawsonia intracellularis growth. Therefore, one other form of this embodiment of the invention relates to vaccines comprising antibodies against the 74 kD Lawsonia intracellularis protein according to 25 the invention.

Vaccines can also be based upon host cells as described above, that comprise the proteins or immunogenic fragments thereof according to the invention.

An alternative and efficient way of vaccination is direct vaccination with DNA encoding the relevant antigen. Direct vaccination with DNA encoding proteins has been successful for many different proteins. (As reviewed in e.g. Donnelly et al., The Immunologist 2: 20-26 (1993)).

This way of vaccination is very attractive for the vaccination of pigs against Lawsonia intracellularis infection.

Therefore, still other forms of this embodiment of the invention relate to vaccines comprising nucleic acids encoding a protein according to the invention or immunogenic fragments thereof according to the invention, and to vaccines comprising DNA fragments that comprise such nucleic acids.

Still other forms of this embodiment relate to vaccines comprising recombinant DNA

DNA vaccines can easily be administered through intradermal application e.g. using a needle-less injector. This way of administration delivers the DNA directly into the cells of the animal to be vaccinated. Amounts of DNA in the microgram range between 1 and 100 µg provide very good results.

molecules according to the invention.

25

In a further embodiment, the vaccine according to the present invention additionally comprises one or more antigens derived from other pig pathogenic organisms and viruses, or genetic information encoding such antigens.

Such organisms and viruses are preferably selected from the group of Pseudorabies virus, Porcine influenza virus, Porcine parvo virus, Transmissible gastro-enteritis virus, Rotavirus, Escherichia coli, Erysipelothrix rhusiopathiae, Bordetella bronchiseptica, Salmonella cholerasuis, Haemophilus parasuis, Pasteurella multocida, Streptococcus suis, Mycoplasma hyopneumoniae, Brachyspira hyodysenteriae and Actinobacillus pleuropneumoniae.

All vaccines according to the present invention comprise a pharmaceutically acceptable carrier. A pharmaceutically acceptable carrier can be e.g. sterile water or a sterile physiological salt solution. In a more complex form the carrier can e.g. be a buffer.

Methods for the preparation of a vaccine comprise the admixing of a protein according to the invention, or an immunogenic fragment thereof, and a pharmaceutically acceptable carrier.

Vaccines according to the present invention may in a preferred presentation also contain an adjuvant. Adjuvants in general comprise substances that boost the immune response of the host in a non-specific manner. A number of different adjuvants are known in the art. Examples of adjuvants are Freunds Complete and Incomplete

adjuvant, vitamin E, non-ionic block polymers, muramyldipeptides, Quill A^(R), mineral oil e.g. Bayol^(R) or Markol^(R), vegetable oil, and Carbopol^(R) (a homopolymer), or Diluvac^(R) Forte.

The vaccine may also comprise a so-called "vehicle". A vehicle is a compound to which the polypeptide adheres, without being covalently bound to it. Often used vehicle compounds are e.g. aluminium hydroxide, -phosphate or -oxide, silica, Kaolin, and Bentonite.

·10

15

20

25

30

A special form of such a vehicle, in which the antigen is partially embedded in the vehicle, is the so-called ISCOM (EP 109.942, EP 180.564, EP 242.380)

In addition, the vaccine may comprise one or more suitable surface-active compounds or emulsifiers, e.g. Span or Tween.

Often, the vaccine is mixed with stabilisers, e.g. to protect degradation-prone polypeptides from being degraded, to enhance the shelf-life of the vaccine, or to improve freeze-drying efficiency. Useful stabilisers are i.a. SPGA (Bovarnik et al; J. Bacteriology 59: 509 (1950)), carbohydrates e.g. sorbitol, mannitol, trehalose, starch, sucrose, dextran or glucose, proteins such as albumin or casein or degradation products thereof, and buffers, such as alkali metal phosphates.

In addition, the vaccine may be suspended in a physiologically acceptable diluent. It goes without saying, that other ways of adjuvating, adding vehicle compounds or diluents, emulsifying or stabilising a polypeptide are also embodied in the present invention.

Vaccines according to the invention can very suitably be administered in amounts ranging between 1 and 100 micrograms, although smaller doses can in principle be used. A dose exceeding 100 micrograms will, although immunologically very suitable, be less attractive for commercial reasons.

Vaccines based upon live attenuated recombinant carriers, such as the LRC-viruses and bacteria described above can be administered in much lower doses, because they

multiply themselves during the infection. Therefore, very suitable amounts would range between 10³ and 10⁹ CFU/PFU for respectively bacteria and viruses.

Many ways of administration can be applied. Oral application is a very attractive way of administration, because the infection is an infection of the digestive tract. A preferred way of oral administration is the packaging of the vaccine in capsules, known and frequently used in the art, that only disintegrate after they have passed the highly acidic environment of the stomach. Also, the vaccine could be mixed with compounds known in the art for temporarily enhancing the pH of the stomach.

Systemic application is also suitable, e.g. by intramuscular application of the vaccine. 10 If this route is followed, standard procedures known in the art for systemic application are well-suited.

From a point of view of protection against disease, a quick and correct diagnosis of Lawsonia intracellularis infection is important.

Therefore it is another objective of this invention to provide diagnostic tools suitable for the detection of Lawsonia intracellularis infection.

A diagnostic test for the detection of Lawsonia intracellularis antibodies in sera can be e.g. a simple standard sandwich-ELISA-test in which 74 kD protein or antigenic 20 fragments thereof according to the invention are coated to the wall of the wells of an ELISA-plate. A method for the detection of such antibodies is e.g. incubation of 74 kD protein or antigenic fragments thereof with serum from mammals to be tested, followed by e.g. incubation with a labelled antibody against the relevant mammalian antibody. A colour reaction can then reveal the presence or absence of antibodies against Lawsonia intracellularis. Another example of a diagnostic test system is e.g. the incubation of a Western blot comprising the 74 kD protein or an antigenic fragment thereof according to the invention, with serum of mammals to be tested, followed by analysis of the blot.

30

25

5

15

Thus, another embodiment of the present invention relates to diagnostic tests for the detection of antibodies against Lawsonia intracellularis. Such tests comprise a protein or a fragment thereof according to the invention.

A diagnostic test based upon the detection of antigenic material of the specific 74 kD protein of Lawsonia intracellularis antigens and therefore suitable for the detection of Lawsonia intracellularis infection can e.g. also be a standard ELISA test. In one example of such a test the walls of the wells of an ELISA plate are coated with antibodies directed against the 74 kD protein. After incubation with the material to be tested, labelled anti-Lawsonia intracellularis antibodies are added to the wells. A colour reaction then reveals the presence of antigenic material from Lawsonia intracellularis.

5

15

20

Therefore, still another embodiment of the present invention relates to diagnostic tests for the detection of antigenic material of Lawsonia intracellularis. Such tests comprise antibodies against a protein or a fragment thereof according to the invention.

The polypeptides or immunogenic fragments thereof according to the invention expressed as characterised above can be used to produce antibodies, which may be polyclonal, monospecific or monoclonal (or derivatives thereof). If polyclonal antibodies are desired, techniques for producing and processing polyclonal sera are well-known in the art (e.g. Mayer and Walter, eds. *Immunochemical Methods in Cell and Molecular Biology*, Academic Press, London, 1987).

Monoclonal antibodies, reactive against the polypeptide according to the invention (or variants or fragments thereof) according to the present invention, can be prepared by immunising inbred mice by techniques also known in the art (Kohler and Milstein, *Nature*, 256, 495-497, 1975).

Methods for large-scale production of antibodies according to the invention are also known in the art. Such methods rely on the cloning of (fragments of) the genetic information encoding the protein according to the invention in a filamentous phage for phage display. Such techniques are described i.a. at the "Antibody Engineering Page" under "filamentous phage display" at http://aximtl.imt.uni-marburg.de/~rek/aepphage.html., and in review papers by Cortese, R. et al., (1994) in Trends Biotechn. 12: 262-267., by Clackson, T. & Wells, J.A. (1994) in Trends Biotechn. 12: 173-183, by Marks, J.D. et al., (1992) in J. Biol. Chem. 267: 16007-16010, by Winter, G. et al., (1994) in Annu. Rev. Immunol. 12: 433-455, and by Little, M. et al., (1994) Biotechn. Adv. 12: 539-555. The phages are subsequently used to screen camelid expression libraries expressing camelid heavy chain

antibodies. (Muyldermans, S. and Lauwereys, M., Journ. Molec. Recogn. 12: 131-140 (1999) and Ghahroudi, M.A. et al., FEBS Letters 414: 512-526 (1997)). Cells from the library that express the desired antibodies can be replicated and subsequently be used for large scale expression of antibodies.

Examples

Example 1:

Isolation of Lawsonia intracellularis from infected porcine ilea.

L. intracellularis infected ilea, confirmed by histopathology and acid-fast Ziehl-5 Neelsen or Whartin-Starry staining, were collected from pigs died with PE, and stored at -80°C. After thawing L. intracellularis bacteria were isolated from mucosal scrapings taken from the infected intestinal wall. The ileal scrapings were homogenized repeatedly in PBS in an omnimixer to release the intracellular bacteria as described by Lawson et al. (Vet. Microbiol. 10: 303-323 (1985)). Supernatant 10 obtained after low-speed centrifugation to remove cell debris was filtered through 5.0, 3.0, 1.2, and 0.8 μm filters (Millipore). The filtrate was subsequently centrifuged at 8000 g for 30 min, giving a small pellet of L. intracellularis bacteria. These bacteria were further purified using a Percoll gradient. The identity of the purified bacteria was assessed by PCR (Jones et al., J. Clin. Microbiol. 31: 2611-2615 (1993)) whereas 15 purity of the isolated bacteria (>95%) was assessed by phase contrast microscopy to reveal any contaminating bacteria or gut debris present.

Bacterial strains and plasmids

L. intracellularis cells were isolated from infected ileal material as described above.
E. coli strain TOP10F' and the TOPO TA cloning kit, containing plasmid pCR2.1
TOPO TA were purchased from Invitrogen (Groningen, the Netherlands). Stocks of all bacterial strains, containing 30% glycerol, were stored at -70°C.
Luria Bertani broth (LB) and LB plates were prepared according to standard
procedures. When needed plasmids were transformed to E. coli TOP10F' competent cells by heat shock. E. coli cells were made competent using standard methods.

DNA isolation

In order to obtain highly purified *L. intracellularis* chromosomal DNA, DNA was prepared from bacterial cells using a Biorad chromosomal DNA isolation kit (Biorad, Veenendaal, the Netherlands) according to manufacturers instructions. Plasmid DNA and linear DNA was isolated using Qiagen products according to the protocols provided by the manufacturer.

PCR amplification

20

25

30

PCR amplification was performed using a Geneamp 9700 PCR system (Applied Biosystems, California, USA). The PCR was performed with the Expand High Fidelity PCR System (Roche Diagnostics GmbH, Mannheim, Germany). The PCR mixture contained 52 U/ml Expand High Fidelity Enzyme Mix, Expand HF buffer with 2.5 mM MgCl₂, 16 mM dNTPs (Promega, Wisconsin, USA), 20 pmoles of primers and 15 ng chromosomal DNA of *L. intracellularis* as template. For standard applications (i.e. colony PCR) the PCR mixture contained 20 U/ml Supertaq and Supertaq buffer (HT Biotechnology Ltd, Cambridge, UK), containing 8 mM dNTPs (Promega, Wisconsin, USA), 10 pmoles of primers and 15 ng template.

In vitro transcription and translation

In vitro transcription and translation was performed using the Rapid Translation

System from Roche Applied Science (Mannheim, FRG) according the manufacturer's protocol.

Summarizing, first the knowledge based sequence-optimization tool ProteoExpert RTS E. coli HY was used to design high yield variants of the original gene. This program optimizes the DNA template for the translation step by suggesting mutations in the DNA sequence. Only silent mutations were allowed, leading to identical aminoacid sequences on the protein level. However, changes of up to 8 nucleotides within the first 6 codons were proposed by the ProteoExpert service to give better expression results.

Ten sense and a universal antisense primers, containing a 5' overlapping region of 20 nucleotides and 15-38 additional gene-specific nucleotides, were used in 10 different PCR reactions to amplify these variants with purified *L. intracellularis* chromosomal DNA as template. The obtained amplicons were purified from gel and used for the generation of linear expression constructs for cell-free protein expression using the RTS E. coli Linear Template Generation Set, His-tag, to introduce the necessary T7 regulatory elements.

Again the obtained amplicons were purified from gel, and after quantification, the appropriate amount of DNA was used for protein expression analysis 50-µl RTS 100 E. coli HY reaction mixture. Expression was analysed using Western blotting with an anti polyhistidine monoclonal antibody.

The construct that gave the highest protein yields was ligated to pCR2.1 TOPO TA vector using the TOPO TA cloning kit. The obtained plasmid was used for medium scale protein production using the RTS 500 E. coli HY kit. The samples were analyzed by SDS page and by Western blot.

The DNA sequence of the expression vector was confirmed using an ABI 310 automated sequencer (Applied Biosystems, California, USA).

Polyacrylamide gel electrophoresis and western blotting

sDS-PAGE was performed using 4-12% Bis-Tris gels from the NuPAGE
electrophoresis system (Novex, San Diego, USA). Western blotting was performed
using semi dry blotting procedures. Western blots were developed using pig antiLawsonia polyclonal serum that was raised against a whole cell preparation in a
water:oil=45:55 emulsion.. The serum was pre-adsorbed using an equal volume crude
cell extracts from BL21star(DE3) containing vector pLysSrare at 4°C for 4 hours.

Results

15

30

Cloning of L. intracellularis gene 5293

For the evaluation of the ProteoExpert suggestions, linear DNA templates were generated via PCR using the RTS Linear Template Generation Set. The primers used in these experiments also introduced a His6-tag at the C-terminus for detection and purification. The PCR-generated templates were examined for their expression performance using RTS 100 E. coli HY Kit. The suggested DNA sequence that gave the highest yields was constructed using primers 5293A5 and 5293B (Table 1) in the first PCR.

The obtained expression construct was ligated to pCR2.1 TOPO TA vector and the resulting vector was transformed to *E. coli* TOP10F and incubated o/n at 37°C. Putative transformants were checked for the right plasmid, using colony PCR. The plasmid inserts, of colony PCR positive transformants, were checked by nucleotide sequence analysis. One of the clones that contained a sequence as expected on basis of the cloning strategy was chosen and designated pTOPO5293.

Table 1. Sequence of the degenerated primers used for the amplification of gene 5293.

Primer	Sequence
5293A5	CTTTAAGAAGGAGATATACCATGGCGGATTATTTAA GTGGTGGAATTTCTTTTGGAGG
5293B	TGATGATGAGAACCCCCCCCCCCCCAAGTTGCC

Expression of L. intracellularis gene 5293 using RTS technology

- Plasmid pTOPO5293 was purified from E. coli TOP10F and the appropriate amount of DNA was added to a RTS500 vial. After incubation conform the protocol of the manufacturer, a sample was taken for analysis using SDS-PAGE gel electrophoresis (Fig. 1A). A clear protein band of approximately 74 kDa was observed in sample that had been taken after 30 hours of induction (Fig. 1A, lane 3) in comparison with the control sample (Fig. 1A, lane 2).
- The same samples were also analysed by western blot using pig serum. The 74 kD protein was specifically recognized by the polyclonal pig serum used in this experiment (Fig 1B, lane 3).
- Conclusion: The 74 kD protein according to the invention can efficiently be expressed and is specifically recognized by the polyclonal pig serum. The 74 kD protein is an important vaccine component for the protection of pigs against Lawsonia intracellularis infection.

Legend to the figure.

Fig. 1. Analysis of the expression of Lawsonia intracellularis gene 5293 using RTS500 technology by SDS-PAGE (A) and Western blotting with polyclonal pig serum (B). Lane 1, molecular weight marker; lane 2, control; lane 3, pET5293 Arrows indicate the location of the expression product.

Claims

- 1) Nucleic acid encoding a 74 kD Lawsonia intracellularis protein or a part of said nucleic acid that encodes an immunogenic fragment of said protein, said nucleic acid or said part thereof having at least 90 %, preferably 92 %, more preferably 94 %, even more preferably 96% homology with a nucleic acid having a sequence as depicted in SEQ ID NO: 1
- 2) DNA fragment comprising a nucleic acid according to claim 1.

5

10

20

- 3) Recombinant DNA molecule comprising a nucleic acid according to claim 1 or a DNA fragment according to claim 2, under the control of a functionally linked promoter.
 - 4) Live recombinant carrier comprising a nucleic acid according to claim 1, a DNA fragment according to claim 2 or a recombinant DNA molecule according to claim 3.
- 15 Host cell comprising a nucleic acid according to claim 1, a DNA fragment according to claim 2, a recombinant DNA molecule according to claim 3 or a live recombinant carrier according to claim 4.
 - 6) A 74 kD Lawsonia intracellularis protein, said protein comprising an amino acid sequence that is at least 90 %, preferably 92 %, more preferably 94 %, even more preferably 96 % homologous to the amino acid sequence as depicted in SEQ ID NO: 2, or an immunogenic fragment of said protein.
 - 7) Lawsonia intracellularis protein according to claim 6 for use in a vaccine.
 - 8) Use of a Lawsonia intracellularis protein according to claim 6 for the manufacturing of a vaccine for combating Lawsonia intracellularis infections.
- 9) Vaccine for combating Lawsonia intracellularis infections, characterised in that it comprises a nucleic acid according to claim 1, a DNA fragment according to claim 2, a recombinant DNA molecule according to claim 3, a live recombinant carrier according to claim 4, a host cell according to claim 5 or a protein according to claim 6, and a pharmaceutically acceptable carrier.
 - 10) Vaccine according to claim 9, characterised in that it comprises an adjuvant.
 - 11) Vaccine according to claim 9 or 10, characterised in that it comprises an additional antigen derived from a virus or micro-organism pathogenic to pigs or genetic information encoding said antigen.

12) Vaccine according to claim 11, characterised in that said virus or microorganism pathogenic to pigs is selected from the group of Pseudorabies virus, Porcine influenza virus, Porcine parvo virus, Transmissible gastro-enteritis virus, Rotavirus, Escherichia coli, Erysipelothrix rhusiopathiae, Bordetella bronchiseptica, Salmonella cholerasuis, Haemophilus parasuis, Pasteurella multocida, Streptococcus suis, Mycoplasma hyopneumoniae, Brachyspira hyodysenteriae and Actinobacillus pleuropneumoniae.

- 13) Vaccine for combating Lawsonia intracellularis infections, characterised in that it comprises antibodies against a protein according to claim 6.
- 14) Method for the preparation of a vaccine according to claim 9-13, said method comprising the admixing of a nucleic acid according to claim 1, a DNA fragment according to claim 2, a recombinant DNA molecule according to claim 3, a live recombinant carrier according to claim 4, a host cell according to claim 5, a protein according to claim 6, or antibodies against a protein according to claim 6, and a pharmaceutically acceptable carrier.
 - 15) Diagnostic test for the detection of antibodies against Lawsonia intracellularis, characterised in that said test comprises a protein or a fragment thereof as defined in claim 6.
 - 16) Diagnostic test for the detection of antigenic material of Lawsonia
 intracellularis, characterised in that said test comprises antibodies against a
 protein or a fragment thereof as defined in claim 6.

Abstract

The present invention relates i.a. to nucleic acids encoding novel Lawsonia intracellularis proteins. It furthermore relates to DNA fragments, recombinant DNA molecules and live recombinant carriers comprising these sequences. Also it relates to host cells comprising such nucleic acids, DNA fragments, recombinant DNA molecules and live recombinant carriers. Moreover, the invention relates to proteins encoded by these nucleotide sequences and to their use for the manufacturing of vaccines. The invention also relates to vaccines for combating Lawsonia intracellularis infections and methods for the preparation thereof.

Finally the invention relates to diagnostic tests for the detection of Lawsonia intracellularis antigens and of antibodies against Lawsonia intracellularis.

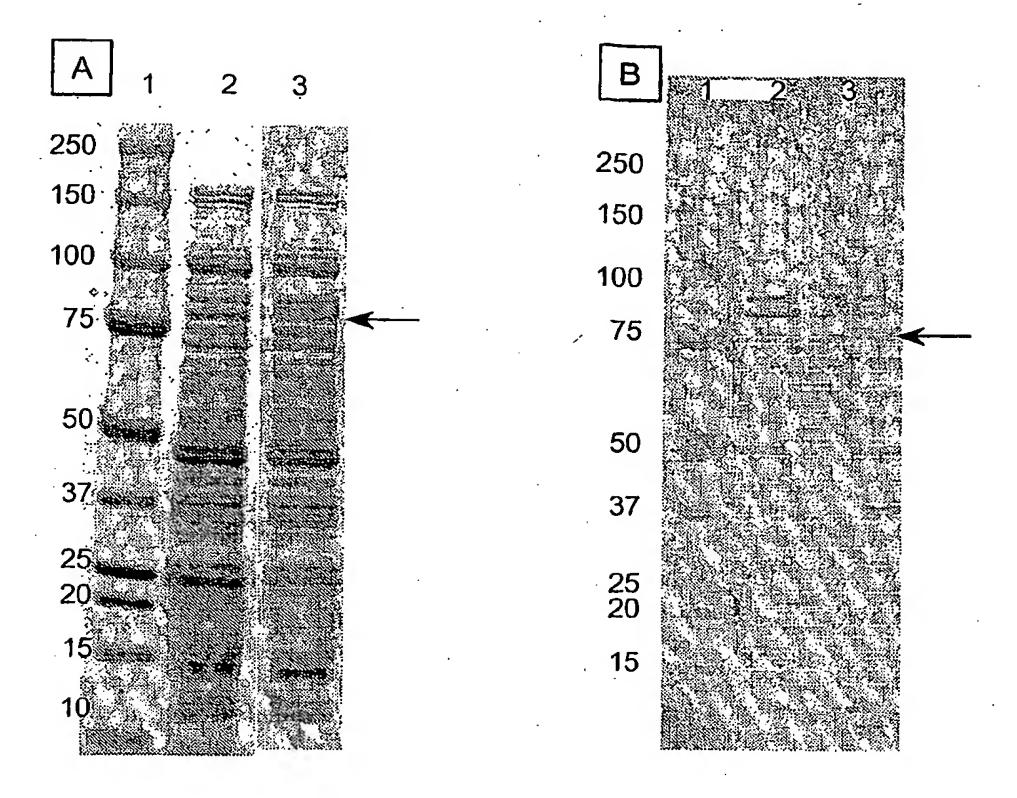


Figure 1.

SEQUENCE LISTING

				•											
<110>	AKZO N	lobel	N.V	7.											·
<120>	Lawson	nia i	ntra	cell	ular	cis 7	74 kI) sub	ounit	vac	cine	:			
<130>	2004.0	06													
<160>	2														
<170>	0> PatentIn version 3.2														
<210> <211> <212> <213>	DNA	nia i	.ntra	acell	lula	ris	·								
<220> <221> <222>	CDS (12).	. (209	96)	•										·	
<400> aggaca	1 aaac t	atg Met 1	gcg	gat Asp	tat	ctt Leu 5	tca Ser	gga Gly	gga Gly	att Ile	tct Ser 10	ttt Phe	gga Gly	gga Gly	50
att gg Ile Gl 15	t agt o	gga a Gly :	acc (Thr)	Asp	ttc Phe 20	caa Gln	gct Ala	atg Met	Ile A	gat Asp 25	caa (Gln)	ctt Leu	aag Lys	aaa Lys	98
att ga 11e G1 30	g ctt u Leu	att (Ile)	Pro	aaa Lys 35	aat Asn	aga Arg	ctt Leu	Val	gtt Val 40	tcc Ser	cat His	gaa Glu	caa Gln	tgg Trp 45	146
aca aa Thr Ly	a aaa ys Lys	Tyr	aaa Lys 50	gca Ala	ttt Phe	gaa Glu	gag Glu	ctt Leu 55	ata Ile	aaa Lys	aca Thr	gtt Val	aaa Lys 60	gat Asp	194
act ga Thr Gl	aa gcg lu Ala	tct Ser 65	tta Leu	agt Ser	aag Lys	cta Leu	agt Ser 70	tct Ser	gtt Val	ggt Gly	gct Ala	att Ile 75	tta Leu	aaa Lys	242
aaa ga Lys Gl	aa ggt Lu Gly 80	tct Ser	gtt Val	tca Ser	aat Asn	act Thr 85	tct Ser	gtt Val	gca Ala	agc Ser	gtt Val 90	aag Lys	gca Ala	agt Ser	290
tct ga Ser As	at gca sp Ala 5	tct Ser	gat Asp	gga Gly	aca Thr 100	cat His	aca Thr	att Ile	gat Asp	gtg Val 105	aaa Lys	cag Gln	ctt Leu	gca Ala	338
aca a: Thr A: 110	ac acg sn Thr	att Ile	ctt Leu	tct Ser 115	aat Asn	aat Asn	cat His	att Ile	ttt Phe 120	gat Asp	tct Ser	aaa Lys	act Thr	gaa Glu 125	386
agt a Ser I	tt aat le Asn	aat Asn	aca Thr 130	ggt Gly	tca Ser	cct Pro	ggt Gly	atc Ile 135	Pne	gct Ala	tat Tyr	gag Glu	tat Tyr 140	77.0	434
GJA G āāā ā	aa cta lu Leu	cat His 145	gaa Glu	gtt Val	gaa Glu	gtt Val	cct Pro 150	Pro	ggt Gly	agt Ser	gat Asp	ctt Leu 155	gaa Glu	tat Tyr	482
ctt g Leu A	ca aca la Thr 160	Leu	ata Ile	aac Asn	aaa Lys	gat Asp 165	Ser	aat Asn	aat Asn	cct Pro	ggt Gly 170	gtt Val	aaa Lys	gca Ala	530
aac c	tt atc	aag	act	ggc	gat	ggc	tat	atg	ttt	agt	ctt	gaa	gga	act	578

	Asn	Leu 175	Ile	Lys	Thr		Asp (180	Gly '	Tyr !	Met	Phe	Ser 185	Leu	Glu	Gly	Thr					
	gaa Glu 190	act Thr	ggt Gly	gca Ala	Asn	gcg Ala 195	act i	tta Leu	tct (Ser	Ile	tca Ser 200	aat Asn	aag Lys	aca Thr	acg Thr	ctt Leu 205	626				
	cca Pro	gac Asp	ttt Phe	Lys	gca Ala 210	tct Ser	gtt Val	gct Ala	Thr	agc Ser 215	agt Ser	gca . Ala	tta Leu	gct Ala	aat Asn 220	ggt Gly	674				
	gaa Glu	gat Asp	aca Thr	att Ile 225	att. Ile	aat Asn	act Thr	Ser	gga Gly 230	aca Thr	act	caa Gln	caa Gln	ttt Phe 235	tct Ser	ttt Phe	722				
	gaa Glu	tạc Tyr	aat Asn 240	gga Gly	aga Arg	aca Thr	Phe	act Thr 245	ttc Phe	gat Asp	att Ile	cct Pro	tca Ser 250	gga Gly	aca Thr	aca Thr	770				
	gca Ala	aaa Lys 255		ctc Leu	caa Gln	aca Thr	gct Ala 260	ata Ile	aat Asn	gaa Glu	aat Asn	aca Thr 265	aaa Lys	aat Asn	aca Thr	Gly	818			•	
	gta Val 270	Arg	gca Ala	act Thr	ttt Phe	gaa Glu 275	Lys	cat His	GJ Y ggc	tca Ser	gat Asp 280	ata Ile	gta Val	ttg Leu	caa Gln	tta Leu 285	866				
	gaa Glu	gga Gly	aca Thr	gtt Val	cct Pro 290	Asn	caa Gln	caa Gln '	gtt Val	aaa Lys 295	val	acc	gct Ala	agc Ser	Pro 300	act Thr	914				
•	gat Asp	ctt Leu	gga Gly	agt Ser 305	Phe	aca Thr	tct Ser	tcg Ser	ggt Gly 310	GIN	gca Ala	Gly	tgg	aat Asn 315	r ny -	e cgt B Arg	962				
	gat Asp	tct Ser	caa Gln 320	Asp	gct Ala	att	ttt Phe	aat Asn 325	тте	aat Asn	ggt Gly	tgg Trp	gac Asp 330	, 611	gaa Glu	e ctt 1 Leu	1010				
	aca Thi	a tct c Sei 335	c Ser	aca Thi	a aat r Ası	gaa n Glu	ctt Lev 340	Tnr	gaa Glu	ı gtt ı Val	ato L Ile	cca Pro 345	, 61	toti Lei	caa 1 Gli	a att n Ile	1058				
·	aca Thi	r Lei	a ctt u Lei	tc: Se:	c gaa r Gl	a ggg u Gly 35	A PAS	a aca 3 Thi	caa Glr	a att	t aca Thi		ca Gl	g ac	t tc r Se	t act r Thr 365	1106				
	ga As	c gaa	a gta u Val	a aa l Ly	a aa s Ly 37	s Gl	a gtt n Val	t gaq 1 Glu	g aaa 1 Lys	a gca s Ala 37	a va.	a gaq 1 Gli	g to 1 Se	t at r Il	a aa e As 38	t aat n Asn 0	1154		-		
	gt Va	t ct 1 Le	t tc u Se.	c aa r Ly 38	s Il	t ca e Gl	a gaq n Glu	g tt: u Le	a actuary This	r rà	a gc	a ac	a gc r Al	t ga a Gl 39	<u>~</u>	c aaa p Lys	1202	•			
	ga As	t ga p As	t ag p Se 40	r Ly	la ga rs As	c ac p Th	t tc r Se	t ag r Se 40	r se	t tc r Se	a ag r Se	t aa r Ly	a at s Il 41	C LL	a to	ea tat er Tyr	1250				
	tt Le	a ca su Gl 41	.n Se	t cc r Pr	et ac	a aa ir Ly	a gt s Va 42	т гл	g gc s Al	t gg a Gl	a ct y Le	a tt u Ph 42	.6 .1.	a gg r Gl	rt ga .y As	at act sp Thr	1298	-			
·	gg G] 43	ly Il	ta ca le Gl	a at n Me	ig ct et Le	et ag eu Se 43	er Th	t ag ır Ar	a ct g Le	t aa eu Ly	ng to ys Se 44	:T T7	c tt e Pi	t to ne Se	et to er Se	et aat er Asn 445	1346		·		
	gģ	gt ct	ta gç	jt t!	tt to	ct co	ct aa	aa ca	a ac	ca ca	aa ga	at go	rt co	ca g	39 g	at cta	139,				,

Gly	Leu	Gly	Phe	Ser 450	Pro	Lys	Gln	Thr	Gln. 455	Asp	Gly	Pro	Gly	Asp 460	Leu	
ttt Phe	tca Ser	tca Ser	ctt Leu 465	gct Ala	tca Ser	att Ile	ggt Gly	att Ile 470	gtc Val	gta Val	gat Asp	gct Ala	gat Asp 475	gag Glu	ggt Gly	1442
agt Ser	gaa Glu	act Thr 480	ttt Phe	gga Gly	caa Gln	ctt Leu	aaa Lys 485	att Ile	tta Leu	gat Asp	aga Arg	gaa Glu 490	aca Thr	att Ile	ggt Gly	1490
cct Pro	gat Asp 495	gca Ala	cct Pro	tat Tyr	aca Thr	act Thr 500	ctt Leu	gat Asp	gag Glu	gca. Ala	tta Leu 505	aaa Lys	aaa Lys	gat Asp	cca Pro	1538
caa Gln 510	gca _. Ala	gta Val	gca Ala	gat Asp	ata Ile 515	tta Leu	gct Ala	ggt Gly	agt Ser	tct Ser 520	gga Gly	ata Ile	tct Ser	gat Asp	tca Ser 525	1586
aca Thr	gat	ttt Phe	tct Ser	tat Tyr 530	caa Gln	gat Asp	cat His	att Ile	gtt Val 535	gga Gly	aaa Lys	aca Thr	caa Gln	gct Ala 540	ggt Gly	1634
Thr	tat Tyr	Asp	Val 545	Lys	Tyr	Ser	Val	Asp 550	Ala	Ser	Gly	Thr	555	СТÀ	qsA	1682
Val	tac	11e 560	GJY	Gly	Val	ГÄ́г	Ala 565	Ser	Leu	Ser	Asp	570	ATa	гла	ASN	1730 /
Ile	tat Tyr 575	Thr	Val	Thr	Ser	Gly 580	Pro	Ala	Thr	Gly	Leu 585	Ser	TTE	Ala	vai	1778
Asn 590	Asn	Arg	Thr	Pro	Gly 595	Ile	Asn	Val	Glu	Ser 600	Thr	Val	Arg	var	603	1826
Gln	Gly	Lys	Leu	Ser 610	Gln	Ile	Gln	Glu	Ala 615	Leu	Lys	Ala	GIU	620		1874
Glr	Asp	Pro	625	Lys	Glu	Asn	Thr	630	Pro	Leu	ıITe	TIE	635	GIII	gat Asp	1922
Ası	тух	640	Asp	Val	Met	Lys	Asn 645	Leu	. Glu	Thr	Arg	650		туу	gaa Glu	
Thi	655	Arç	y Val	. Thr	: Ser	660	Glu	· Arg	, wet	; met	665	; Tet	ı pås	File	tct Ser	
aga Are 67	g Let	gat Asp	gct Ala	gta Val	tta Lev 675	Ala	l aaa Lys	tat Tyr	aat Asr	caç Glr 680	n Met	g ato : Met	g tca Ser	gca Ala	a aat a Asn 685	
gc Al	t tct a Sei	agt c Sei	t tta c Lev	ggg 1 Gly 690	, Glr	ctt Let	ggt Gly	gea Ala	a taa	a						2096

<210> 2 <211> 694

<211> 694 <212> PRT

<213> Lawsonia intracellularis

<400> 2

Met Ala Asp Tyr Leu Ser Gly Gly Ile Ser Phe Gly Gly Ile Gly Ser 1 5 10 15

Gly Thr Asp Phe Gln Ala Met Ile Asp Gln Leu Lys Lys Ile Glu Leu 20 25 30

Ile Pro Lys Asn Arg Leu Val Val Ser His Glu Gln Trp Thr Lys Lys 35 40 45

Tyr Lys Ala Phe Glu Glu Leu Ile Lys Thr Val Lys Asp Thr Glu Ala 50 55 60

Ser Leu Ser Lys Leu Ser Ser Val Gly Ala Ile Leu Lys Lys Glu Gly 65 70 75 80

Ser Val Ser Asn Thr Ser Val Ala Ser Val Lys Ala Ser Ser Asp Ala 85 90 95

Ser Asp Gly Thr His Thr Ile Asp Val Lys Gln Leu Ala Thr Asn Thr 100 105 110

Ile Leu Ser Asn Asn His Ile Phe Asp Ser Lys Thr Glu Ser Ile Asn 115 120 125

Asn Thr Gly Ser Pro Gly Ile Phe Ala Tyr Glu Tyr Lys Gly Glu Leu 130 135 140

His Glu Val Glu Val Pro Pro Gly Ser Asp Leu Glu Tyr Leu Ala Thr 145 150 150

Leu Ile Asn Lys Asp Ser Asn Asn Pro Gly Val Lys Ala Asn Leu Ile 165 170 175

Lys Thr Gly Asp Gly Tyr Met Phe Ser Leu Glu Gly Thr Glu Thr Gly 180

Ala Asn Ala Thr Leu Ser Ile Ser Asn Lys Thr Thr Leu Pro Asp Phe 195 200 205

Lys Ala Ser Val Ala Thr Ser Ser Ala Leu Ala Asn Gly Glu Asp Thr 210 215 220

Ile Ile Asn Thr Ser Gly Thr Thr Gln Gln Phe Ser Phe Glu Tyr Asn 235 240

Gly Arg Thr Phe Thr Phe Asp Ile Pro Ser Gly Thr Thr Ala Lys Glu 245

Leu Gln Thr Ala Ile Asn Glu Asn Thr Lys Asn Thr Gly Val Arg Ala

260 265 270

Thr Phe Glu Lys His Gly Ser Asp Ile Val Leu Gln Leu Glu Gly Thr 275 280 285

Val Pro Asn Gln Gln Val Lys Val Thr Ala Ser Pro Thr Asp Leu Gly 290 295 300

Ser Phe Thr Ser Ser Gly Gln Ala Gly Trp Asn Lys Arg Asp Ser Gln 305 310 315

Asp Ala Ile Phe Asn Ile Asn Gly Trp Asp Gln Glu Leu Thr Ser Ser 325

Thr Asn Glu Leu Thr Glu Val Ile Pro Gly Leu Gln Ile Thr Leu Leu 340 345 350

Ser Glu Gly Lys Thr Gln Ile Thr Ile Gln Thr Ser Thr Asp Glu Val 355 360 365

Lys Lys Gln Val Glu Lys Ala Val Glu Ser Ile Asn Asn Val Leu Ser 370 380

Lys Ile Gln Glu Leu Thr Lys Ala Thr Ala Glu Asp Lys Asp Asp Ser 390 395 400

Lys Asp Thr Ser Ser Ser Ser Lys Ile Pro Ser Tyr Leu Gln Ser 405

Pro Thr Lys Val Lys Ala Gly Leu Phe Thr Gly Asp Thr Gly Ile Gln 420 425 430

Met Leu Ser Thr Arg Leu Lys Ser Ile Phe Ser Ser Asn Gly Leu Gly 435

Phe Ser Pro Lys Gln Thr Gln Asp Gly Pro Gly Asp Leu Phe Ser Ser 450

Leu Ala Ser Ile Gly Ile Val Val Asp Ala Asp Glu Gly Ser Glu Thr 480

Phe Gly Gln Leu Lys Ile Leu Asp Arg Glu Thr Ile Gly Pro Asp Ala 485

Pro Tyr Thr Thr Leu Asp Glu Ala Leu Lys Lys Asp Pro Gln Ala Val

Ala Asp Ile Leu Ala Gly Ser Ser Gly Ile Ser Asp Ser Thr Asp Phe 515

Ser Tyr Gln Asp His Ile Val Gly Lys Thr Gln Ala Gly Thr Tyr Asp

530 535 540

Val Lys Tyr Ser Val Asp Ala Ser Gly Thr Ile Gly Asp Val Tyr Ile 545 550 550

Gly Gly Val Lys Ala Ser Leu Ser Asp Pro Ala Lys Asn Ile Tyr Thr 565 570 575

Val Thr Ser Gly Pro Ala Thr Gly Leu Ser Ile Ala Val Asn Asn Arg 580 585 590

Thr Pro Gly Ile Asn Val Glu Ser Thr Val Arg Val Lys Gln Gly Lys 595 600 605

Leu Ser Gln Ile Gln Glu Ala Leu Lys Ala Glu Val Gln Gln Asp Pro 610 620

Leu Lys Glu Asn Thr Gly Pro Leu Ile Ile Met Gln Asp Asn Tyr Lys 635 630 635

Asp Val Met Lys Asn Leu Glu Thr Arg Ile Glu Lys Glu Thr Gln Arg 645 650 655

Val Thr Ser Trp Glu Arg Met Met Arg Leu Lys Phe Ser Arg Leu Asp 660 665 670

Ala Val Leu Ala Lys Tyr Asn Gln Met Met Ser Ala Asn Ala Ser Ser 675 680 685

Leu Gly Gln Leu Gly Ala 690